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09/932,474	08/17/2001	Timothy E. Benson	00236.U\$1	6629
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MUETING, RAASCH & GEBHARDT, P.A. P.O. BOX 581415			STEADMAN, DAVID J	
MINNEAPOLIS, MN 55458		ART UNIT	PAPER NUMBER	
			1652	
			DATE MAILED: 10/01/2004	

Please find below and/or attached an Office communication concerning this application or proceeding.

AND THE RESIDENCE OF THE PROPERTY OF THE PROPE		T 2				
	Application No.	Applicant(s)				
	09/932,474	BENSON ET AL.				
Office Action Summary	Examiner	Art Unit				
	David J Steadman	1652				
The MAILING DATE of this communication ap	opears on the cover sheet with the o	correspondence address				
A SHORTENED STATUTORY PERIOD FOR REPITHE MAILING DATE OF THIS COMMUNICATION  - Extensions of time may be available under the provisions of 37 CFR 1 after SIX (6) MONTHS from the mailing date of this communication.  - If the period for reply specified above is less than thirty (30) days, a re  - If NO period for reply is specified above, the maximum statutory period  - Failure to reply within the set or extended period for reply will, by statu Any reply received by the Office later than three months after the mailine earned patent term adjustment. See 37 CFR 1.704(b).		mely filed  ys will be considered timely. In the mailing date of this communication.  ED (35 U.S.C. § 133).				
Status						
1) Responsive to communication(s) filed on 28.	Julv 2004.					
·= · ·	·					
3) Since this application is in condition for allow						
closed in accordance with the practice under	closed in accordance with the practice under Ex parte Quayle, 1935 C.D. 11, 453 O.G. 213.					
Disposition of Claims						
4)  Claim(s) 1-57 is/are pending in the applicatio 4a) Of the above claim(s) 7-50 is/are withdray 5)  Claim(s) is/are allowed. 6)  Claim(s) 1-6 and 51-57 is/are rejected. 7)  Claim(s) is/are objected to. 8)  Claim(s) are subject to restriction and/	vn from consideration.					
Application Papers						
9)⊠ The specification is objected to by the Examin	ner.					
10) ☐ The drawing(s) filed on 17 August 2001 is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.						
Applicant may not request that any objection to the						
Replacement drawing sheet(s) including the corre						
Priority under 35 U.S.C. § 119						
<ul> <li>12) Acknowledgment is made of a claim for foreign a) All b) Some * c) None of:</li> <li>1. Certified copies of the priority document 2. Certified copies of the priority document 3. Copies of the certified copies of the priority application from the International Bureat * See the attached detailed Office action for a list</li> </ul>	nts have been received. nts have been received in Applicat ority documents have been receive au (PCT Rule 17.2(a)).	ion No ed in this National Stage				
Attachment(s)						
1) Notice of References Cited (PTO-892)	4) Interview Summary					
<ul> <li>2) Notice of Draftsperson's Patent Drawing Review (PTO-948)</li> <li>3) Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08 Paper No(s)/Mail Date 3/25/02; 3/11/03.</li> </ul>	Paper No(s)/Mail D					

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### **DETAILED ACTION**

# Status of the Application

[1] Claims 1-57 are pending in the application.

### Election/Restriction

- [2] Applicants' election without traverse of Group I, claims 1-6 and 51-57, filed July 28, 2004, is acknowledged.
- [3] Claims 7-50 are withdrawn from further consideration pursuant to 37 CFR 1.142(b) as being drawn to a nonelected invention, there being no allowable generic or linking claim. Election was made without traverse in the reply filed on July 28, 2004.

## **Priority**

[4] Applicant's claim for domestic priority under 35 USC § 119(e) to provisional application numbers 60/226,239 and 60/226,269, both filed August 17, 2000, is acknowledged.

## Information Disclosure Statement

[5] All references cited in the information disclosure statements (IDS) filed March 25, 2002 have been considered by the examiner. All references with the exception of "Database Accession No." references cited in the IDS filed March 11, 2003 have been considered by the examiner. "Database Accession No." references have not been

considered as there is no citation of the relevant database and there is no date of publication. A copy of each IDS is attached to the instant Office action.

## Specification/Informalities

- The attempt to incorporate subject matter into this application by reference to a hyperlink embedded in the specification, *e.g.*, page 31, line 12 and all other hyperlinks embedded in the specification, is improper. Incorporation of subject matter into the patent application by reference to a hyperlink and/or other forms of browser-executable code is considered to be an improper incorporation by reference. See MPEP 608.01 regarding hyperlinks in the specification and 608.01(p), paragraph I regarding incorporation by reference.
- [7] The specification is objected to as referring to "Table 1" (e.g., page 4, line 9), however, there does not appear to be a Table 1 in the specification. Applicants are requested to correct this discrepancy.

# Claim Rejections - 35 USC § 112, Second Paragraph

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

[8] Claim(s) 1-6 and 51-57 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

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- [a] Claim 1, 2 (claim 3 dependent therefrom), 4 (claim 5 dependent therefrom), and 6 are indefinite in the recitation of "amino acids listed in Table..." Without a reference amino acid sequence, e.g., SEQ ID NO:1, it is unclear as to the relative positions of amino acids listed in the recited Table in the molecule or molecular complex. It is suggested that applicants clarify the meaning of the claims.
- [b] Claim 6 is indefinite in the recitation of "structurally homologous to an *S. aureus* FemA molecule or molecular complex" as it is unclear as to whether the recited "structure" is intended to be interpreted as the primary amino acid sequence or the three-dimensional structure. Further, it is unclear as to how "homologous" a molecule or molecular complex must be to be included within the scope of the claim. Also, it is unclear as to the proteins that are considered to be *S. aureus* FemA molecules or molecular complexes such that a skilled artisan can determine whether the claimed molecule or molecular complex is sufficiently "structurally homologous" to be included within the scope of the claim. It is suggested that applicants clarify the meaning of the term.
- [c] Claim 6 is indefinite in the recitation of "represented by" as it is unclear as to the intended meaning of the term in the context of a molecule or molecular complex. It is suggested that applicants clarify the meaning of the term.
- [d] Claim 51-55 are indefinite in the recitation of "S. aureus FemA" as it is unclear as to the scope of proteins that are meant to be encompassed by the terms. It is suggested that applicants clarify the meaning of the claims by providing distinguishing characteristics of an "S. aureus FemA" protein such that a skilled artisan can distinguish

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those proteins that are intended to be encompassed within the scope of the claims from those that are not. For example, by identifying the "S. aureus FemA" by a sequence identifier.

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- [e] Claim 56 is unclear in the recitation of "including amino acids having the sequence of SEQ ID NO:1." It is unclear from the claim and the specification as to whether the phrase is meant to be interpreted as meaning the crystallized *S. aureus* FemA polypeptide has the sequence of SEQ ID NO:1, or if the crystal comprises an *S. aureus* FemA polypeptide and further includes "amino acids having the sequence SEQ ID NO:1." In the interest of advancing prosecution, the phrase has been interpreted as meaning the crystallized *S. aureus* FemA polypeptide has the sequence of SEQ ID NO:1. It is suggested that applicants clarify the meaning of the claim.
- Claims 56 and 57 are confusing in that the claims are drawn to a crystal of an *S. aureus* FemA polypeptide of SEQ ID NO:1 or a selenomethionine variant of SEQ ID NO:1. However, the prior art (Tschierske et al. *FEMS Microbiol Lett* 171:97-102; cited in the IDS filed March 25, 2002) teaches that a polypeptide that is 100% identical to SEQ ID NO:1, referred to as "FmhC," has only 37% amino acid sequence identity to *S. aureus* FemA (page 99, Table I). Thus, it is unclear as to how the polypeptide of SEQ ID NO:1 is considered to be an *S. aureus* FemA. Applicants are requested to clarify this discrepancy.

Claim Rejections - 35 USC § 101

35 U.S.C. 101 reads as follows:

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Whoever invents or discovers any new and useful process, machine, manufacture, or composition of matter, or any new and useful improvement thereof, may obtain a patent therefor, subject to the conditions and requirements of this title.

[9] Claim(s) 1-6 are rejected under 35 U.S.C. 101 because the claimed invention is directed to non-statutory subject matter. The claim(s) are drawn to a molecule or molecular complex. The claim(s) read on a product of nature and should be amended to indicate the hand of the inventor, e.g., by insertion of "purified" or "isolated". See MPEP § 2105.

# Claim Rejections - 35 USC § 112, First Paragraph

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

[10] Claim(s) 1-6 and 51-57 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention. Claims 1-6 are drawn to a genus of molecules or molecular complexes comprising at least a portion of an *S. aureus* FemA or FemA-like substrate binding surface or a genus of molecules or molecular complexes that are homologous thereto. Claim 51 is drawn to a method for crystallizing a genus of *S. aureus* FemA molecules or molecular complexes. Claims 52-57 are drawn to a genus of *S. aureus* FemA crystals.

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For claims drawn to a genus, MPEP § 2163 states the written description requirement for a claimed genus may be satisfied through sufficient description of a representative number of species by actual reduction to practice, reduction to drawings, or by disclosure of relevant, identifying characteristics, i.e., structure or other physical and/or chemical properties, by functional characteristics coupled with a known or disclosed correlation between function and structure, or by a combination of such identifying characteristics, sufficient to show the applicant was in possession of the claimed genus. See Eli Lilly, 119 F.3d at 1568, 43 USPQ2d at 1406. MPEP § 2163 states that a "representative number of species" means that the species which are adequately described are representative of the entire genus. Thus, when there is substantial variation within the genus, one must describe a sufficient variety of species to reflect the variation within the genus. In this case, the specification discloses only a single representative species of molecules or molecular complexes comprising at least a portion of an S. aureus FemA or FemA-like substrate binding surface or a genus of molecules or molecular complexes that are homologous thereto, i.e., the S. aureus FemA of SEQ ID NO:1 and the specification discloses only a single representative species of S. aureus FemA crystals, i.e., a crystal of the purified S. aureus FemA of SEQ ID NO:1 having an orthorhombic space group symmetry of P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> and the unit cell dimensions of a=53.9Å, b=90.4Å, and c=109.3 Å and  $\alpha$ = $\beta$ = $\gamma$ =90° (see particularly p. 15 of the specification). The specification fails to describe any additional representative species of the recited genus of molecules or molecular complexes comprising at least a portion of an S. aureus FemA or FemA-like substrate binding

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surface or *S. aureus* FemA crystals as encompassed by the claims. In this case, the single representative species fails to describe each genus of molecules or molecular complexes or crystals, which encompasses *widely* variant species with respect to the structures and functions of the molecules or molecular complexes and the *S. aureus* FemA protein sequences and crystal structures of the *S. aureus* FemA crystals. Given the lack of description of a representative number of species, the specification fails to sufficiently describe the claimed invention in such full, clear, concise, and exact terms that a skilled artisan would recognize that applicant was in possession of the claimed invention.

[11] Claims 1-6 and 51-57 are rejected under 35 U.S.C. 112, first paragraph, because the specification, while being enabling for the *S. aureus* FemA polypeptide of SEQ ID NO:1; a method of crystallizing the *S. aureus* FemA polypeptide of SEQ ID NO:1 by preparing purified *S. aureus* FemA polypeptide of SEQ ID NO:1 to a final concentration of 12 mg/mL and crystallizing *S. aureus* FemA polypeptide of SEQ ID NO:1 by hanging drop method in a solution of 4% PEG 8000, 100 mM Tris buffer, pH 8.5 at 4 degrees Celsius, a method of crystallizing the *S. aureus* FemA polypeptide of SEQ ID NO:1 having methionine replaced with selenomethionine by preparing purified selenomethionine *S. aureus* FemA polypeptide of SEQ ID NO:1 to a final concentration of 12 mg/mL and crystallizing selenomethionine *S. aureus* FemA polypeptide of SEQ ID NO:1 by hanging drop method in a solution of 30% PEG 4000, 100 mM Tris buffer, pH 8.5, 200 mM MgCl<sub>2</sub> at 20 degrees Celsius, and a crystal of the purified *S. aureus* FemA of SEQ ID NO:1 having an orthorhombic space group symmetry of P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> and the unit

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cell dimensions of a=53.9Å , b=90.4Å , and c=109.3 Å and  $\alpha$ = $\beta$ = $\gamma$ =90°, does not reasonably provide enablement for all molecules or molecular complexes as encompassed by claims 1-6, methods for crystallizing any *S. aureus* FemA polypeptide under the broad range of conditions as recited in claim 51, or all crystals of an *S. aureus* FemA polypeptide as encompassed by claims 52-57. The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make the invention commensurate in scope with these claims.

It is the examiner's position that undue experimentation would be required for a skilled artisan to make the entire scope of the claimed invention. Factors to be considered in determining whether undue experimentation is required are summarized in *In re Wands* (858 F.2d 731, 737, 8 USPQ2d 1400, 1404 (Fed. Cir. 1988)) as follows: (A) The breadth of the claims; (B) The nature of the invention; (C) The state of the prior art; (D) The level of one of ordinary skill; (E) The level of predictability in the art; (F) The amount of direction provided by the inventor; (G) The existence of working examples; and (H) The quantity of experimentation needed to make or use the invention based on the content of the disclosure. See MPEP § 2164.01(a). The Factors most relevant to the instant rejection are addressed in detail below.

• The claims are overly broad in scope: Claims 1-6 are so broad as to encompass all molecules or molecular complexes comprising at least a portion of an *S. aureus*FemA substrate binding surface or site as encompassed by claims 1-5 or structural homologs thereof as encompassed by claim 6. As claims 1-6 have been given their

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broadest reasonable interpretation in accordance with MPEP 2111, the claimed molecules and molecular complexes encompass essentially any protein. Claim 51 is so broad as to encompass a method for crystallizing any S. aureus FemA polypeptide. including all mutants and variants of the polypeptide of SEQ ID NO:1. Further, it is noted that the claim recites an extremely broad set of crystallization conditions. Claims 52-56 are so broad as to encompass a vast number of S. aureus FemA crystals, having any space group symmetry and/or unit cell dimensions, including all mutants and variants of the polypeptide of SEQ ID NO:1. In this case the disclosure is limited to the S. aureus FemA polypeptide of SEQ ID NO:1; a method of crystallizing the S. aureus FemA polypeptide of SEQ ID NO:1 by preparing purified S. aureus FemA polypeptide of SEQ ID NO:1 to a final concentration of 12 mg/mL and crystallizing S. aureus FemA polypeptide of SEQ ID NO:1 by hanging drop method in a solution of 4% PEG 8000. 100 mM Tris buffer, pH 8.5 at 4 degrees Celsius, a method of crystallizing the S. aureus FemA polypeptide of SEQ ID NO:1 having methionine replaced with selenomethionine by preparing purified selenomethionine S. aureus FemA polypeptide of SEQ ID NO:1 to a final concentration of 12 mg/mL and crystallizing selenomethionine S. aureus FemA polypeptide of SEQ ID NO:1 by hanging drop method in a solution of 30% PEG 4000, 100 mM Tris buffer, pH 8.5, 200 mM MgCl<sub>2</sub> at 20 degrees Celsius, and a crystal of the purified S. aureus FemA of SEQ ID NO:1 having an orthorhombic space group symmetry of P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> and the unit cell dimensions of a=53.9Å, b=90.4Å, and c=109.3

Å and  $\alpha = \beta = \gamma = 90^{\circ}$ .

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The lack of guidance and working examples: The specification provides only a single working example of the claimed molecule or molecular complex, i.e., the S. aureus FemA polypeptide of SEQ ID NO:1. The specification provides only two working examples of the claimed method of crystallization, i.e., a method of crystallizing the S. aureus FemA polypeptide of SEQ ID NO:1 by preparing purified S. aureus FemA polypeptide of SEQ ID NO:1 to a final concentration of 12 mg/mL and crystallizing S. aureus FemA polypeptide of SEQ ID NO:1 by hanging drop method in a solution of 4% PEG 8000, 100 mM Tris buffer, pH 8.5 at 4 degrees Celsius, a method of crystallizing the S. aureus FemA polypeptide of SEQ ID NO:1 having methionine replaced with selenomethionine by preparing purified selenomethionine S. aureus FemA polypeptide of SEQ ID NO:1 to a final concentration of 12 mg/mL and crystallizing selenomethionine S. aureus FemA polypeptide of SEQ ID NO:1 by hanging drop method in a solution of 30% PEG 4000, 100 mM Tris buffer, pH 8.5, 200 mM MgCl<sub>2</sub> at 20 degrees Celsius. The specification provides only a single working example of the claimed crystal, i.e., a crystal of the purified S. aureus FemA of SEQ ID NO:1 having an orthorhombic space group symmetry of P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> and the unit cell dimensions of a=53.9Å, b=90.4Å, and c=109.3 Å and  $\alpha = \beta = \gamma = 90^{\circ}$ . These working examples fail to provide the necessary guidance for making the entire scope of crystal compositions broadly encompassed by the claims. Regarding claims 1-6, the specification fails to provide guidance regarding those amino acids of SEQ ID NO:1 that may be altered by substitution, addition, insertion, and/or deletion with an expectation of maintaining the desired FemA activity.

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Furthermore, the specification fails to provide guidance as to how to use those variant polypeptides having activities other than the desired FemA activity, *e.g.*, non-functional polypeptides or polypeptides having activity other than FemA activity. Regarding claims 51-57, the specification fails to provide guidance regarding crystallization of other *S. aureus* FemA proteins using other methods/conditions of crystallization with an expectation of obtaining diffraction quality crystals.

• The high degree of unpredictability in the art is supported by the state of the art: Regarding claims 1-6, the amino acid sequence of a protein determines its structural and functional properties. Predictability of which changes can be tolerated in a protein's amino acid sequence and obtain the desired FemA activity requires a knowledge of and guidance with regard to which amino acids in the protein's sequence, if any, are tolerant of modification and which are conserved (i.e., expectedly intolerant to modification), and detailed knowledge of the ways in which the proteins' structure relates to its function. The positions within a protein's sequence where modifications can be made with a reasonable expectation of success in obtaining a polypeptide having the desired FemA activity/utility are limited and the result of such modifications is highly unpredictable. In addition, one skilled in the art would expect any tolerance to modification for a given protein to diminish with each further and additional modification, e.g., multiple substitutions.

The state of the art provides evidence for the high degree of unpredictability in altering a protein's sequence with an expectation that the encoded polypeptide will maintain the desired activity/utility. For example, Branden et al. ("Introduction to Protein

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Structure", Garland Publishing Inc., New York, 1991) teach "[p]rotein engineers frequently have been surprised by the range of effects caused by single mutations that they hoped would change only one specific and simple property in enzymes" and "[t]he often surprising results of such experiments reveal how little we know about the rules of protein stability... ... they also serve to emphasize how difficult it is to design *de novo* stable proteins with specific functions" (page 247). The teachings of Branden et al. are exemplified by Witkowski et al. (*Biochemistry* 38:11643-11650), who teach that a single amino acid substitution results in conversion of the parent polypeptide's activity from a beta-ketoacyl synthase to a malonyl decarboxylase (see e.g., Table 1, page 11647). Thus, the prior art acknowledges the unpredictability of altering a protein sequence with an expectation of obtaining a protein having a desired function and discloses that even a single substitution in a polypeptide's amino acid sequence may completely alter the function of a polypeptide.

Regarding claims 51-57, Branden et al. ("Introduction to Protein Structure Second Edition", Garland Publishing Inc., New York, 1999) teach that protein crystallization is usually quite difficult to achieve and the formation of protein crystals is critically dependent on a number of different parameters, including pH, temperature, protein concentration, the nature of the solvent and precipitant, as well as the presence of added ions and ligands to the protein (page 375, middle). Branden et al. teach that even small changes in the crystallization parameters, e.g., pH, can cause the molecules to pack in different ways to produce different crystal forms (page 375, bottom). Thus, even minor modifications to a crystallization method may result in crystals that are

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distinct in structure having different space group symmetry and unit cell dimensions. At least in view of the teachings of Branden et al. a skilled artisan would recognize the high degree of unpredictability in generating the broad scope of claimed crystals.

This unpredictability is evidenced by the teachings of the specification, which describes the difficulties of protein concentration/solubilization (pp. 47-48) and the difficulty of obtaining diffraction-quality crystals of *S. aureus* FemA of SEQ ID NO:1 at 20 degrees Celsius.

• The amount of experimentation required is undue: Regarding claims 1-6, while methods of generating variants of a given protein are known in the art, e.g.,, site-directed mutagenesis, it is not routine in the art to screen for all proteins having a substantial number of substitutions or modifications and having any function, as encompassed by the instant claims. Regarding claims 51-57, while methods of protein crystallization are known, it is not routine in the art to screen a vast number proteins under any crystallization conditions or to experiment to make all crystals as broadly encompassed by the claims.

In view of the overly broad scope of the claims, the lack of guidance and working examples provided in the specification, the high degree of unpredictability as evidenced by the prior art, and the amount of experimentation required to make and use all molecules or molecular complexes, make all methods, or make all crystals as broadly encompassed by the claims, undue experimentation would be necessary for a skilled artisan to make and use the entire scope of the claimed invention.

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Thus, applicant has not provided sufficient guidance to enable one of ordinary skill in the art to make and use the claimed invention in a manner reasonably correlated with the scope of the claims. The scope of the claims must bear a reasonable correlation with the scope of enablement (*In re Fisher*, 166 USPQ 19 24 (CCPA 1970)). Without sufficient guidance, determination of having the desired biological characteristics is unpredictable and the experimentation left to those skilled in the art is unnecessarily, and improperly, extensive and undue. See *In re Wands* 858 F.2d 731, 8 USPQ2nd 1400 (Fed. Cir, 1988).

# Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

- (b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.
- [12] Claim(s) 1-6 are rejected under 35 U.S.C. 102(b) as being anticipated by Tschierske et al. (*FEMS Microbiol Lett* 171:97-102; cited in the IDS filed March 25, 2002) as evidenced by GenBank Accession Number AF106851.

The claims are drawn to a molecule or molecular complex comprising at least a portion of an *S. aureus* FemA or FemA-like substrate binding surface or site or a molecular complex homologous thereto.

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Tschierske et al. teaches an *S. aureus* polypeptide referred to as "FmhC" and teach the sequence of FmhC has GenBank Accession Number AF106851 (page 99, bottom). GenBank Accession Number AF106851 teaches a polypeptide that is 100% identical to SEQ ID NO:1 of the instant application (see Appendices A and B). This anticipates claims 1-6 as written.

#### Citation of Relevant Art

[13] The following reference is made of record as the reference is considered to be pertinent to applicant's disclosure: Benson et al. *Structure* 10:1107-1115. The reference is not relied upon for a prior art rejection as the reference was published after the effective filing date of the instant application.

#### Conclusion

- [14] Status of the claims:
- Claims 1-57 are pending.
- Claims 7-50 are withdrawn from further consideration.
- Claims 1-6 and 51-57 are rejected.
- No claim is in condition for allowance.
- Claims 51-57 would appear to be allowable if rewritten or amended to overcome the rejection(s) under 35 U.S.C. 112, 2nd paragraph, set forth in this Office action.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to David Steadman, whose telephone number is (571) 272-0942. The Examiner can normally be reached Monday-Friday from 6:30 am to 4:00 pm. If attempts to reach the Examiner by telephone are unsuccessful, the Examiner's

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supervisor, Ponnathapura Achutamurthy, can be reached at (571) 272-0928. The FAX

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number for submission of official papers to Group 1600 is (703) 872-9306. Draft or informal FAX communications should be directed to (571) 273-0942. Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the Art Unit receptionist whose telephone number is (703) 308-0196.

David J. Steadman, Ph.D.

**Primary Examiner** 

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09-28-04